



# Why Do Agroforestry Systems Enhance Biodiversity? Evidence From Habitat Amount Hypothesis Predictions

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Considering the present ecological crisis, land use-biodiversity relationships have become a major topic in landscape planning, ecosystem management and ecological restoration. In this scope, consistent patterns of outstanding biodiversity have been identified in agroforestry systems within diverse biogeographic regions and types of management. Empirical work has revealed that agroforestry higher structural complexity, when compared with current simplified agricultural systems, might be partially responsible for the observed patterns. The recently developed Habitat Amount Hypothesis predicts diversity for a local habitat patch, from the amount of the same habitat within the local landscape. We have expanded the previous hypothesis to the landscape level, computing the influence of the dominant land uses on the diversity of coexisting guilds. As a case study, we have considered archetypal landscapes dominated (or co-dominated) by crops or trees, which were compared using normalized diversities. The results obtained show that agroforestry systems substantially increase functional diversity and overall biodiversity within landscapes. We highlight that the normalized values should be parametrized to real conditions where the type of crop, tree and agroecological management will make a difference. Most importantly, our findings provide additional evidence that agroforestry has a critical role in enhancing biodiversity in agricultural landscapes and, in this way, should be regarded as a priority measure in European Agri-environmental funding schemes.

**Keywords:** land-sharing, land-sparing, agroecosystems, trade-offs, landscape heterogeneity, fragmentation, biodiversity management

## INTRODUCTION

Farming systems should be recognized not only from the crops' production perspective, but also from the regulating and cultural services standpoint (Plieninger et al., 2019). In fact, several works highlight their contribution to biodiversity conservation, soil enrichment, landscape beauty and carbon sequestration (e.g., Santos M. et al., 2019; Liu et al., 2020). However, the adoption of intensive management techniques, new crops, large-scale machinery and the massive use of fertilizers and pesticides triggered obvious changes in farming systems and agricultural landscapes, with significant drawbacks (Kanter et al., 2018; Bakış et al., 2021). In Europe, the Common Agricultural Policy (CAP) encouraged the specialization of agriculture and forestry systems in the most productive areas and abandonment of marginal (less productive) regions (De Roest et al., 2018; Santos et al., 2018; Abson, 2019). This was especially noticed in southern Europe, where CAP supporting schemes (including Agri-environmental measures) promoted relevant socio-ecological change but were unable to halt biodiversity loss and stop ecosystem services' degradation (Harlio et al., 2019; Pardo et al., 2020).

Agroforestry, a jeopardized land use system in which trees are grown in combination with crops and/or livestock systems, is considered as one of the fundamental strategies to tackle the increasing need for high quality productions while maximizing ecosystem services and reducing environmental impacts (Torralba et al., 2016; Arosa et al., 2017; Den Herder et al., 2017; de Jalón et al., 2018; Moreno et al., 2018). In fact, wood pastures and grazed orchards, which are still common landscape features in some parts of Mediterranean Europe (De Roest et al., 2018), were highlighted as win-win multifunctional systems, responding to consumers' worries while embracing the united nations sustainable development goals (van Noordwijk et al., 2018). Furthermore, multifunctional benefits of introducing trees in arable lands, novel silvopastoralism techniques and the maintenance and/or improvement of already existing agroforestry practices were recently discussed and added to the European Union Green Deal<sup>1</sup>.

Agroforestry has been identified as important for reducing species loss in agricultural landscapes, but also for endangered species conservation (e.g., Mosquera-Losada et al., 2009; Torralba et al., 2016; Udawatta et al., 2019). Biodiversity conservation is one of Europe 2030 objectives for a "resource efficient" Europe, contributing to a range of services that benefit human wellbeing, including food and fiber production and regulating and cultural services (EU Biodiversity Strategy for 2030)<sup>2</sup>. Nevertheless, species loss has been particularly dramatic in the Mediterranean (namely in agroecosystems), considered a biodiversity hotspot (Palacín and Alonso, 2018; Rosas-Ramos et al., 2019). In this way, understanding how agroforestry influences patterns of biodiversity is of utmost importance, given its significance for future landscape planning and for funding the most effective Agri-environment measures (Ansell et al., 2016;

Pavlis et al., 2016). Agroforestry has the potential to contribute to biodiversity conservation in agricultural landscapes (and forest landscapes) by increasing structural complexity, and enhancing habitat and landscape heterogeneity (Torralba et al., 2016; Boinot et al., 2019; Hagggar et al., 2019). Nevertheless, limited understanding of the interrelated effects of agroforestry micro-habitats on diversity patterns reduces our ability to project how a possible expansion of agroforestry will impact biodiversity and ecosystem services (Boinot et al., 2019; Santos P.Z.F. et al., 2019; Richard et al., 2020).

Biodiversity patterns and strategies for biodiversity conservation in agricultural landscapes have been increasingly enlightened by the recently developed Habitat Amount Hypothesis (HAH) (e.g., Fahrig, 2013, 2021; Melo et al., 2017; Watling et al., 2020). The HAH suggests that species richness, occurrence and abundance (from here referred to as "diversity") in a given habitat site depend exclusively on the amount of that habitat in the "local landscape", defined for an area surrounding the site (effect of scale) (Watling et al., 2020). Even though some authors challenged the HAH, considering "Spatial Configuration" or "Island Effect" theories more important to describe and manage biodiversity in agricultural landscapes (Hanski, 2015; Evju and Sverdrup-Thygeson, 2016; Haddad et al., 2017), new research shows that HAH is actually complementary and might integrate the previous concepts (Bueno and Peres, 2019; Watling et al., 2020; Fahrig, 2021; Saura, 2021). Indeed, Saura (2021) suggested that most discussion might have aroused from a misinterpretation of the HAH, i.e., habitat/landscape configuration is actually fundamental for HAH biodiversity patterns. Also, and as noticed in the original paper, HAH is not the only determinant of diversity: other factors should be considered, such as the type of species vs the type of habitat, the appropriate scale to analyze and the quality of the habitat matrix (Fahrig, 2013).

The practical implementation of the HAH was explicitly depicted by Saura (2021), assuming a formula that relates diversity in the habitat site with the amount of the same habitat within the local landscape. Saura (2021) does not investigate the different types of "diversities" that are associated with the different types of habitats encompassed by a landscape. This is especially pertinent when estimating diversity in complex landscapes or within the intermingled micro-habitats of agroforestry (Riginos et al., 2009; Arosa et al., 2017; Simonson et al., 2018). To decode this gap, a gradient of landscapes should be tested (e.g., dominated by agroforestry, agriculture, forest), estimating the potential diversity provided by each land-use type. Also, as the type of species (e.g., guilds such as habitat specialists vs edge species) is of utmost importance when applying the HAH, this was also simulated in our investigation, since the interaction between type of species and their habitat might change in accordance with the environmental conditions (Watling et al., 2020).

A promising way to gain insight of how HAH predicts diversity patterns in agricultural landscapes is to create virtual landscapes using spatial models (Mas et al., 2014), which enable the integration of habitat specific information and estimation of emergence outcomes (e.g., Prevedello et al.,

<sup>1</sup>[https://www.europarl.europa.eu/RegData/etudes/BRIE/2020/651982/EPRS\\_BRI\(2020\)651982\\_EN.pdf](https://www.europarl.europa.eu/RegData/etudes/BRIE/2020/651982/EPRS_BRI(2020)651982_EN.pdf)

<sup>2</sup>[https://ec.europa.eu/environment/nature/biodiversity/strategy/index\\_en.htm](https://ec.europa.eu/environment/nature/biodiversity/strategy/index_en.htm)

2016; Santos et al., 2016). Also, simple spatial explicit models can improve our understanding of diversity within ecological systems and support the identification of landscape thresholds and successful conservation practices (Sequeira et al., 2018; Bakış et al., 2021). In the present work, a simple and flexible two-dimensional spatial model was developed, focusing on the interaction between diversity and landscapes, easily adaptable to other regions and case studies. In fact, identifying how different agricultural landscapes are related with biodiversity (and concomitant ecosystem services) is fundamental to assess whether agroforestry provides a biodiverse (and productive) alternative to contemporary Mediterranean agricultural landscapes (Kay et al., 2019a).

The following hypotheses were tested: (1) HAH should predict diversity at each site using the landscape matrix; (2) HAH discriminates the different guilds associated with each site; (3) HAH landscape diversity emerges from the combined effects of the size and isolation of all patches in the landscape; (4) HAH anticipates the creation of highly diverse agricultural landscapes when agroforestry is the dominant land use.

## METHODS

### Virtual Landscape Description

The virtual landscape mimics characteristic agricultural landscapes of Mediterranean Europe (Debussche et al., 1999; Van Doorn and Bakker, 2007; Geri et al., 2010): agricultural landscapes dominated by crops (from here referred to as “Agriculture”); agricultural landscapes dominated by trees (from here referred to as “Forest”); agricultural landscapes integrating in separate areas crops and trees (from here referred to as “Clumped”); agricultural landscapes alternating rows of crops with rows of trees (modern agroforestry, from here referred to as “Linear”); agricultural landscapes integrating random patches of crops and trees (traditional agroforestry, from here referred to as “Random”) (Figure 1). The conceptual description of the model follows a simplified version of the specifications of the standard protocol ODD (Overview, Design Concepts and Details) (Grimm et al., 2010). The software Netlogo 6.1.1 (Wilensky, 1999) was chosen to create virtual landscapes and to calculate the biodiversity outcomes.

### Purpose

The model investigates how agricultural landscapes biodiversity emerges from the interplay between patch-level land uses, discriminated within tree or crop dominated patches. By implementing the HAH (Watling et al., 2020) to the simulated landscapes, we intend to understand agroforestry biodiversity patterns, compare with other agricultural landscapes related to agricultural intensification and/or abandonment/afforestation (Santos et al., 2016).

### Entities, State Variables and Scales

The model includes 10,000 patches (unit cells) that make up the landscape extent of the study area (Figure 1 and Table 1 for details). Each patch was characterized by its dominant land

use (tree or crop) and the normalized biodiversity was gauged accordingly with the formulas of Saura (2021), applied to all patches of the landscape.

## Agricultural Landscape Simulations

Simple agricultural landscapes presented only crop patches (Agriculture) or tree patches (Forest) while in the case of Linear, Random and Clumped agricultural landscapes, an equivalent number of patches is associated with crops and trees (Figure 1).

## Diversity Predictions

Diversity predictions ( $D_i$ ) per patch were partitioned within crop species biodiversity ( $C_B$ ), tree species biodiversity ( $T_B$ ), edge species biodiversity ( $E_B$ ) and overall species biodiversity ( $B$ , sums the contributions of the previous guilds) (e.g., Morrison et al., 2017; Wilson et al., 2017; Jackson et al., 2019).  $C_B$  and  $T_B$  were assessed for crop and tree patches, respectively.  $E_B$  was gauged only when the number of dissimilar bordering patches was equal or greater than 1. The power function was used to relate Diversity ( $D_i$ ) in a specific patch, considering the habitat amount surrounding it within a radius of  $i$  ( $A_i$ ) (total area of habitat matching the central patch), as defined by Saura (2021) (Eq. 1). The power function associated with the HAH was calibrated to follow an average slope ( $z$ ) of 0.45, supported on previous studies (Haddad et al., 2017; Rabelo et al., 2017; Bueno and Peres, 2019; Saura, 2021). In addition,  $k$  is a constant which depends on the unit used for area measurement, and equals the diversity that would exist if the habitat area was confined to one square unit (Whittaker et al., 2017) (details in Supplementary Appendix 5).

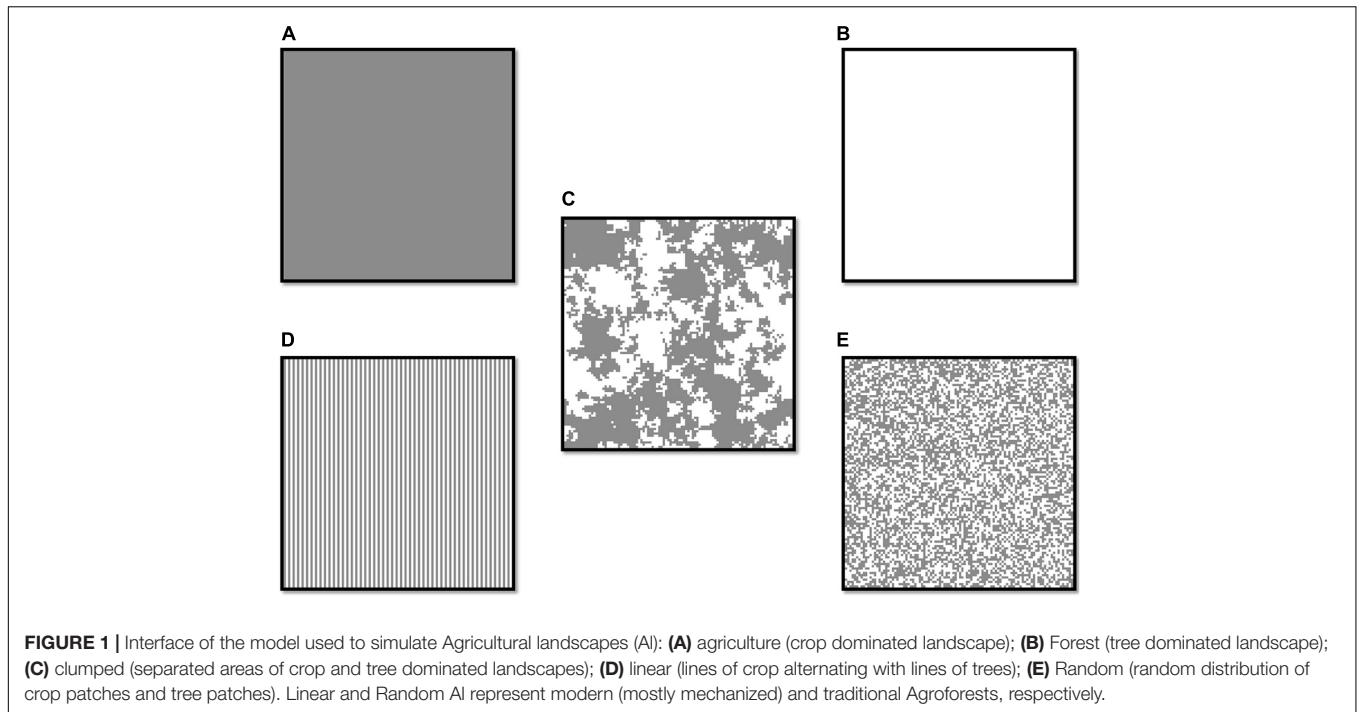
$$D_i = k * A_i^{0.45} \quad (1)$$

All metrics of diversity ( $C_B$ ,  $T_B$ ,  $E_B$ ,  $B$ ) were normalized ( $D_{norm,i}$ , Eq. 2) using  $D_{max}$ , the maximum value of  $D_i$  possible.  $D_{norm,i}$  ( $N_C_B$ ,  $N_T_B$ ,  $N_E_B$ ,  $N_B$ ) values, ranging from 0 to 1, are independent from  $k$  and, in this way, from the type of diversity, thus allowing direct comparisons between the different metrics and the different landscapes (Geneletti et al., 2018; Saura, 2021) (details in Supplementary Appendix 5).

$$D_{norm,i} = \frac{D_i}{D_{max}} \quad (2)$$

## Comparing the Agricultural Landscapes' Diversities

Cohen's effect size was used to reveal the magnitude of the differences in diversity between the agricultural landscapes simulated (Sullivan and Feinn, 2012; White et al., 2014). The rules of thumb for standardized mean differences effect sizes are defined as:  $d = 0$ , none;  $d \approx 0.01$ , very small;  $d \approx 0.2$ , small;  $d \approx 0.5$ , medium;  $d \approx 0.8$ , large;  $d \approx 1.2$ , very large;  $d \approx (>) 2$ , huge (Sawilowsky, 2009). To complement the previous results, the Kruskal-Wallis test and the Steel-Dwass multi-comparisons test were applied to determine the significance of pairwise differences (Sokal and Rohlf, 1995; Zar, 1996). Statistics were done using R 4.0.3 (R Core Team, 2020), through the effectsize (v 0.4.0; Ben-Shachar et al., 2020) and the ezr packages (Kanda, 2013).



## RESULTS

Habitat Amount Hypothesis biodiversity predictions were uneven, contingent on the guild composition and agricultural landscape (**Figure 2**). In fact, while Normalized crop biodiversity ( $N_{C_B}$ ) and Normalized tree biodiversity ( $N_{T_B}$ ) attained higher medians in Agriculture and Forest dominated landscapes, respectively, Normalized Edge Biodiversity ( $N_{E_B}$ ) and Normalized Biodiversity ( $N_B$ ) reached higher medians within Agroforestry landscapes (Linear and Random) (**Figure 2**). On the other hand, the lower medians for  $N_{C_B}$ ,  $N_{T_B}$  and  $N_{E_B}$  were simulated for Forest, Agriculture and both Agricultural landscapes, respectively (**Figure 2**). Additionally, Clumped landscapes presented the higher dispersion (interquartile range) for all guilds ( $N_{C_B}$ ,  $N_{T_B}$ ,  $N_{E_B}$ ,  $N_B$ ) (**Figure 2**). Moreover, all Biodiversity was condensed within a single partition in Agriculture and Forest landscapes,  $N_{C_B}$  and  $N_{F_B}$ , respectively, and spread within all guilds ( $N_{C_B}$ ,  $N_{T_B}$ ,  $N_{E_B}$ ) for Clumped, Linear and Random landscapes (**Figures 2A–C**).

Pairwise differences using effect sizes ( $d$ ) corroborate the previous results, depicting that even though Clumped landscapes, incorporating both uses (crop and tree dominated patches) in separate areas, are expected to be associated with more biodiversity ( $N_B$ ) than single Agriculture or Forest (Very small differences,  $d < 0.2$ ), only Agroforestry (Linear and Random landscapes) differences (from Agriculture and Forest) were considered Huge ( $d > 2.0$ ) (**Table 2**). Additionally, both types of Agroforestry depict none to small differences in the biodiversity metrics ( $N_{C_B}$ ,  $N_{T_B}$ ,  $N_{E_B}$ ), although with a small advantage in overall biodiversity for linear Agroforestry ( $N_B$ ,  $d = 0.249$ ) (**Table 2**).

Complementary comparisons using non-parametric statistics, depicted in the **Supplementary Material (Supplementary Appendixes 1–4)**, corroborate the previous results:  $N_{C_B}$  higher  $t$ -values were attained when comparing Agriculture with all other agricultural landscapes (**Supplementary Appendix 1**);  $N_{T_B}$  higher  $t$ -values were attained when comparing Forest with all other agricultural landscapes (**Supplementary Appendix 2**);  $N_{E_B}$  higher  $t$ -values were attained when comparing Agroforestry landscapes (Linear and Random) with all other agricultural landscapes (**Supplementary Appendix 3**);  $N_B$  higher  $t$ -values were attained when comparing Agroforestry landscapes (Linear and Random) with all other agricultural landscapes, particularly Linear vs Agriculture, Forestry and Clumped (**Supplementary Appendix 4**).

## DISCUSSION

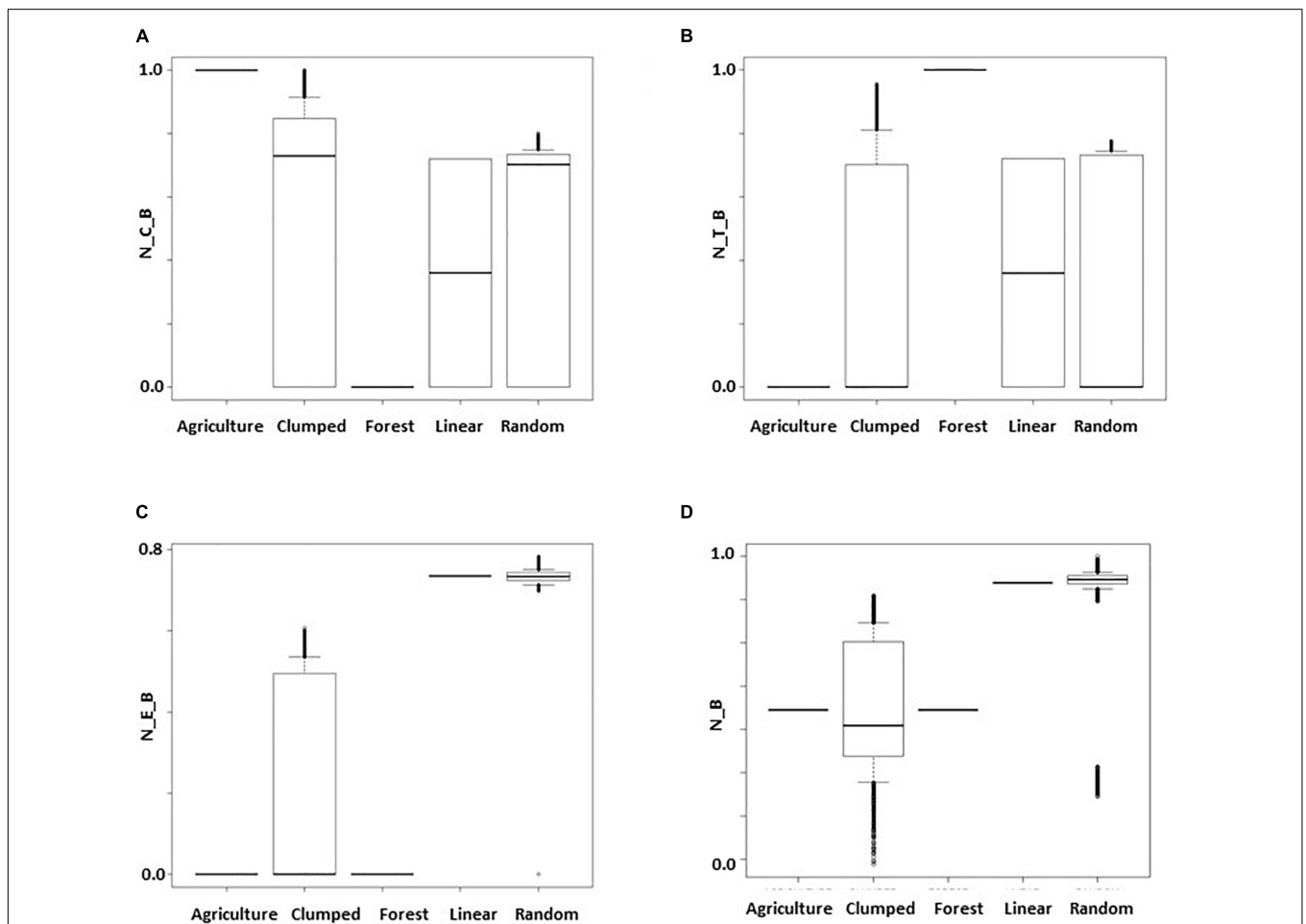
### Agricultural Landscapes' Complexity and Biodiversity Estimates

As expected (e.g., Fahrig, 2013; Torralba et al., 2016), higher Normalized crop biodiversity ( $N_{C_B}$ ) and Normalized tree biodiversity ( $N_{T_B}$ ) values were attained within monospecific landscapes, Agriculture and Forest correspondingly, but also both landscapes low overall biodiversity' predictions (Normalized biodiversity,  $N_B$ ) (**Figure 2** and **Table 2**). The corresponding higher dispersion for all guilds in the case of Clumped landscapes [ $N_{C_B}$ ,  $N_{T_B}$ ,  $N_{E_B}$  (Normalized edge biodiversity)] was also anticipated (**Figure 2**): depending of the fractal structure of the landscape associated with the type of species and grain size, detached patches with very low biodiversity could occur close to biodiverse ones (e.g., Benton et al., 2003; Toderi et al., 2017;



**TABLE 1** | Description of the variables of each conceptual entity used in the model.

Entity	Variable	Brief description
Patches	Number	– 10,000 patches arranged in a square grid (100 × 100 patches), which corresponds to a total of 100 ha.
	Size	– Individual patch – 100 square meters (10 × 10 meters).
Land-use-Diversity		– Each patch corresponds to a single dominant land use, crop or tree, identified by different colors.
		– Depending on the type of Land-use, biodiversity (Crop biodiversity, C_B and Tree biodiversity, T_B) is gauged considering the habitat area surrounding within a radius of $i$ ( $A_i$ ), using Eq. 1 (see please Methods – Diversity predictions). In the case of Edge biodiversity (E_B), the calculations are made considering 4 as the maximum number of edges by patch (four possible neighbors with different land uses).
N_Diversity		– Biodiversity was normalized using Eq. 2.
Landscape		– Landscape sets the patches dispersion/distribution, from uniform, to linear, passing through random or clumped. The agricultural landscapes considered were: (a) dominated by crops, Agriculture; (b) integrating in separate areas crops and trees, Clumped; (c) dominated by trees, Forest; (d) alternating rows of crops with rows of trees, Linear; (e) integrating random patches of crops and trees, Random. Linear and Random are associated with Agroforestry.



**FIGURE 2** | Box and Whisker plots expressing the differences in the Biodiversity metrics associated with the different Agricultural landscapes simulated: **(A)** normalized Crop Biodiversity (N\_C\_B); **(B)** normalized Tree Biodiversity (N\_T\_B); **(C)** normalized Edge Biodiversity (N\_E\_B); **(D)** normalized Biodiversity (N\_B). Medians for N\_C\_B: Agriculture, 1.000; Clumped, 0.728; Forest, 0.000; Linear, 0.360; Random, 0.702. Medians for N\_T\_B: Agriculture, 0.000; Clumped, 0.000; Forest, 1.000; Linear, 0.360; Random, 0.000. Medians for N\_E\_B: Agriculture, 0.000; Clumped, 0.000; Forest, 0.000; Linear, 0.735; Random, 0.735. Medians for N\_B: Agriculture, 0.645; Clumped, 0.609; Forest, 0.645; Linear, 0.938; Random, 0.945. Associated Steel-Dwass multiple comparisons tests in **Supplementary Appendixes 1–4**.

Bakış et al., 2021). Nevertheless, even if Agroforestry landscapes (Linear and Random) were associated with comparable N\_C\_B and N\_T\_B to Clumped landscapes, their maximization of ecotones had an overwhelming impact on the N\_E\_B and in the

N\_B (Figure 2C and Table 2) as suspected from previous works (Jose, 2012; Moreno et al., 2018; Marconi and Armengot, 2020). This last comparison was surprising as, from the HAH theory predictions, equivalent amounts of habitat (Agroforestry – Linear

and Random – had equivalent areas per habitat to Clumped) should have resulted in equivalent diversity (Fahrig, 2013, 2021; Saura, 2021).

The results obtained when applying the HAH to our virtual agricultural landscapes contribute to pinpoint the general mechanisms underlying biodiversity enhancement by agroforestry systems (Udawatta et al., 2019). In particular, agroforestry “boosting” effect was captured by the HAH implementation, since HAH implicitly reflects isolation/linkage of habitat patches and complexity within landscapes which is inherently associated with the biodiversity predictions (Bueno and Peres, 2019; Balzan et al., 2020; Saura, 2021). Also, the outcomes we got agree with Torralba et al. (2016) meta-analysis of European agroforestry: the authors found that land-use systems that are structurally and functionally more complex than crop-dominated or tree-dominated monospecific systems result in higher levels of biodiversity and ecosystem goods and services, without compromising productivity (see also Jose, 2012; Haggard et al., 2019).

## Issues of Scale, Management and Integration With Metapopulation Dynamics

Our results support the hypothesis that agroforestry impacts biodiversity (Torralba et al., 2016). However, our simple model does not consider mechanisms operating at larger scales, neither management issues or crop/tree species’ combinations (Hanane et al., 2019; Kay et al., 2019b; Varah et al., 2020). Habitat heterogeneity at different scales – from individual farms to whole landscapes – is commonly recognized as the core factor to reverse declines in farmland biodiversity (Benton et al., 2003). Albeit current Agri–environment measures to sustain biodiversity being implemented at the farm level, their ultimate objective is to impact landscapes (Toderi et al., 2017). Yet, the impact of the resulting landscape-scale distribution of habitat patches and their effectiveness remains largely unknown (Ansell et al., 2016; Marja et al., 2019). Ultimately, agroforestry benefits to the society depend on upscaling the different practices to generate add-value at landscape scale, particularly if applied in the right locations and with the correct management options (Plieninger et al., 2020).

Management plays an essential role shaping agroforestry biodiversity: positive effects of unproductive and reduced input areas and negative effects of intensive management were noticed for diverse taxonomic groups and systems (Lüscher et al., 2016). As an example, the shift to organic farming combined with agroforestry can increase biodiversity further, not only agrobiodiversity (e.g., traditional varieties) but also wild species associated with the complex microenvironments created (e.g., Marconi and Armengot, 2020; Rosati et al., 2020). Also, environmentally friendly agroforestry management might improve valuable ecosystem services such as habitat for pollinators and natural enemies of pests (Rosati et al., 2020). However, despite the diverse Agri–environmental measures of the Common Agricultural Policy (CAP), European agriculture is ruled by productivity, aggravating biodiversity loss (Concepción et al., 2020). Actually, agroforestry benefits

for biodiversity – and agroforestry long term sustainability – might be jeopardized when these systems are oversimplified (e.g., Steffan-Dewenter et al., 2007; Arosa et al., 2017; Avilés, 2019).

The crop(s) and tree(s) chosen have an enormous effect on biodiversity enhancement in agroforestry, namely when native and/or traditional crops are combined with native tree species (Campos et al., 2007; Torralba et al., 2016). Native trees, apart from the structural complexity gain associated with agroforestry, support higher diversity of resources for wildlife, are usually adapted to local conditions and might even contribute to reduce fire risk in Mediterranean ecosystems (Damianidis et al., 2020). Conversely, alien species, especially potentially invasive trees, might threaten biodiversity and should be totally avoided (McNeely, 2004). Nevertheless, most productive non-invasive alien tree species could be considered, if applying adaptive management and monitoring, namely in harsh environmental conditions (e.g., eroded soils) (Guillerme et al., 2020).

Additionally, even if the HAH predictions seem to integrate the metapopulation dynamics theory (Semper-Pascual et al., 2021), by holistically considering that extinction probability decreases as the size of patches of habitat increase (assuming that expected population size is positively correlated with overall habitat area), the distributions of organisms do not always reflect the distribution of suitable habitats (Merckx et al., 2019). Population dynamics were not explicitly considered in our simple model, nor changes in quality of the micro-habitats present (Gardiner et al., 2018). Nevertheless, several authors consider that more attention should be given to spatially explicit methods for hierarchically defining patches based on the dynamics of the focal species (e.g., Cavanaugh et al., 2014). Moreover, patch occupancy, colonization and extinction result from combination of processes at both local and regional scales, especially difficult to be understood and simulated for multispecies responses to landscape dynamics (Dallas et al., 2020).

## From the Patch to the Landscape: Expanding the Habitat Amount Hypothesis

Biodiversity predictions from HAH should be analyzed not only as a whole but considering the different partitions (e.g., guilds), assuming that mechanisms of biodiversity emerge from patch-scale changes, such as edge effects, changes in behavior and local interactions (Fletcher et al., 2018). The underlying mechanistic pathway occurs *via* expanding connectivity and dispersal, while suitability for each species might vary spatially within patches and in relation to configuration variables (e.g., distance from edge) (Fletcher et al., 2018). The approach used in this work was also able to highlight that even though agroforestry should increase overall landscape biodiversity, part of this diversity – associated with crops and or trees – is only attained when the previous land uses dominate the landscape (Sokos et al., 2013; Concepción and Díaz, 2019). That is to say that agroforestry may not be a panacea for halting biodiversity loss in agroecosystems but, within an increasingly anthropogenic ecosystems and novel landscapes’ world, one of the win-win strategies (Montagnini, 2017). In fact, the relevance for overall biodiversity of tree patches (and

**TABLE 2 |** Pairwise comparisons (Cohen's *d*) for the Normalized diversity predictions between all Agriculture landscapes (Agricultural landscapes details in **Table 1**): interpretation of Cohen's *d* values and average *d* values in parenthesis. Positive values indicate higher diversity for the first landscape in comparison, negative values indicate higher diversity for the second landscape in comparison, and 0 indicates equivalent diversity values. N\_C\_B, normalized crop biodiversity; N\_T\_B, normalized tree biodiversity; N\_E\_B, normalized edge biodiversity; N\_B, normalized biodiversity. Complementary Kruskal-Wallis and Stoll-Dwass multiple comparisons test in **Supplementary Appendixes 1–4**.

Pairwise comparisons	N_C_B	N_T_B	N_E_B	N_B
<i>Agriculture vs Clumped</i>	Very large (1.771)	Very large (−1.162)	Very large (−1.306)	Very small (−0.099)
<i>Agriculture vs Forest</i>	Huge (1e + 07)	Huge (−1e + 06)	None (0.000)	None (0.000)
<i>Agriculture vs Linear</i>	Huge (2.514)	Very large (−1.414)	Huge (−7.3e + 06)	Huge (−2.9 e + 06)
<i>Agriculture vs Random</i>	Huge (2.419)	Very large (−1.402)	Huge (−5.618)	Huge (−3.455)
<i>Clumped vs Forest</i>	Very large 1.671	Huge (−2.709)	Very large (1.306)	Very small (0.099)
<i>Clumped vs Linear</i>	Small (0.324)	Small (−0.165)	Huge (−2.869)	Huge (−2.793)
<i>Clumped vs Random</i>	Small (0.294)	Small (−0.170)	Huge (−2.149)	Huge (−2.048)
<i>Forest vs Linear</i>	Very large (−1.414)	Huge (2.514)	Huge (−7.3e + 06)	Huge (−2.9 e + 06)
<i>Forest vs Random</i>	Very large (−1.425)	Huge (2.469)	Huge (−5.618)	Huge (−3.455)
<i>Linear vs Random</i>	Very small (−0.029)	None (−0.006)	Small (0.353)	Small (0.249)

isolated trees) in agroecosystems was highlighted by Concepción et al. (2020) but also the detrimental effects on agroecosystem specialists (e.g., threaten pseudo-steppe species).

The HAH challenges the assumption that small patches have little biodiversity value, by predicting that the size of the patch in which a plot is located has little additional effect on diversity beyond its contribution to the habitat amount in the local landscape (Saura, 2021). Also, evidence shows that the effect of habitat amount around a sample plot is stronger than either the individual or combined effects of patch size and isolation (Watling et al., 2020). In our case study, landscape biodiversity emerges from the several patches specific contributions to habitat amount. When the local landscape is composed by a single use, overall landscape biodiversity decreases, even though with increasing values for specific guilds (e.g., N\_C\_B or N\_T\_B). We have assumed that the larger the area occupied by a single land use, more important is its contribution to increase habitat amount, but not more than a collection of multiple patches summing an equivalent area. Nevertheless, higher biodiversity is expected to occur in more complex landscapes (e.g., comparing Clumped vs Random or Linear landscapes), suggesting agroforestry as a welcomed strategy in conservation.

## Final Remarks Concerning Food Production, Agroforestry Systems and Sustainability

Higher crop yields and food production needed for an increasingly population, originated an intense debate whether the most assertive strategy to biodiversity conservation is land-sharing (low-yield, environmentally friendly agriculture occupying larger areas, represented here by agroforestry) or land-sparing (high-yield, conventionally intensified agriculture occupying smaller areas, represented by Clumped) (e.g., Grass et al., 2019). Our conceptual model seems to support the idea that intensification may foster biodiversity declines at the landscape scale, as well as at the farm site through the direct effect of homogenization (e.g., Phalan et al., 2011). Thus, land-sharing might be the winning strategy for biodiversity

enhancement, without disregarding the importance of land-sparing for specialists and endangered species conservation (Marull et al., 2018). In fact, multi-scale perspectives are fundamental and should consider land use history, local values, soil characteristics, climate but also socio-economic factors influence on land use at the regional scale and external drivers such as agricultural policies, trade treaties or climate agreements (Karner et al., 2019).

Mediterranean agriculture urges alternatives and agroforestry could be a key element amongst the tools to fight contemporary environmental challenges, such as climate change, water scarcity and food security (Malek et al., 2018; Yves et al., 2020). Concerning biodiversity, trees in agricultural landscapes appear particularly efficient in contributing to biodiversity conservation, while environmentally valuable and economically profitable (Barrios et al., 2018; Kay et al., 2019b). In the European Union, the “green architecture” of the new CAP framework (2021–2028) includes the safeguarding of small landscape elements and connectivity features, as well as regionally-adapted and specific voluntary environmental measures (such as agroforestry) (Nilsson et al., 2017; Ustaoglu and Collier, 2018; Balzan et al., 2020; Hristov et al., 2020). Ultimately, agroforestry systems address several sustainable development goals, offer countless ecosystem services and (hopefully) are expected to get more attention and importance in the future of world's agriculture (Kanter et al., 2018).

## CONCLUSION

Our framework complements previous approaches relating agroforestry and biodiversity conservation in agricultural landscapes. The simple modeling demonstration, supported on a robust theory, could be considered a stepping-stone to support future quantification of biodiversity patterns within agricultural landscapes. Further unraveling of how habitat amount influences biodiversity in agricultural landscapes should consider other variables such as type of species (e.g., guilds),

management practices and multiple spatial and temporal scales. Our findings also reinforce the idea that agroforestry is critical to halt biodiversity loss in agricultural landscapes and, despite the limitations inherent to a preliminary demonstration, the methodology developed provides a starting point to anticipate the changes in landscape biodiversity induced by land use change and management options, guiding pertinent strategies to integrate crop production with biodiversity conservation.

## DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding author.

## AUTHOR CONTRIBUTIONS

MS: conceptualization, formal analysis, investigation, software, writing – original draft, and writing – review and editing. RC: conceptualization and writing – review and editing. RB, DG, A-LP, PL, WB, and JC: writing – review and editing. DG: writing – review and editing. BG and MM-L: conceptualization and writing – review and editing. All authors contributed to the article and approved the submitted version.

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## SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fevo.2021.630151/full#supplementary-material>

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